

- (1) GENERAL INFORMATION:
 - (i) APPLICANT: Sidransky, David
 - (ii) TITLE OF INVENTION: NUCLEIC ACID MUTATION DETECTION IN HISTOLOGIC TISSUE
 - (iii) NUMBER OF SEQUENCES: 82
 - (iv) CORRESPONDENCE ADDRESS:
 - (A) ADDRESSEE: Spensley Horn Jubas & Lubitz(B) STREET: 1880 Century Park East, Suite 500
 - (C) CITY: Los Angeles
 - (D) STATE: California
 - (E) COUNTRY: USA (F) ZIP: 90067
 - (v) COMPUTER READABLE FORM:

 - (A) MEDIUM TYPE: Floppy disk (B) COMPUTER: IBM PC compatible (C) OPERATING SYSTEM: PC-DOS/MS-DOS
 - (D) SOFTWARE: PatentIn Release #1.0, Version #1.25
 - (vi) CURRENT APPLICATION DATA:
 - (A) APPLICATION NUMBER: US
 - (B) FILING DATE: JANUARY 14, 1994
 - (C) CLASSIFICATION:

 - (viii) ATTORNEY/AGENT INFORMATION:
 (A) NAME: Wetherell, Jr., Ph.D., John R.
 (B) REGISTRATION NUMBER: 31,678
 (C) REFERENCE/DOCKET NUMBER: PD-3055

 - (ix) TELECOMMUNICATION INFORMATION:
 - (A) TELEPHONE: (619) 455-5100 (B) TELEFAX: (619) 455-5110
- (2) INFORMATION FOR SEQ ID NO:1:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 26 base pairs
 - (B) TYPE: nucleic acid (C) STRANDEDNESS: single

 - (D) TOPOLOGY: linear
 - (ii) MOLECULE TYPE: DNA (genomic)
 - (ix) FEATURE:
 - (A) NAME/KEY: CDS (B) LOCATION: 1..26
 - (xi) SEQUENCE DESCRIPTION: SEQ ID NO:1:

AAGTCAGGGC ACAAGTGAAT TCCTAC

- (2) INFORMATION FOR SEQ ID NO:2:
 - (i) SEQUENCE CHARACTERISTICS:

	(i)	SEQUENCE CHARACTERISTICS: (A) LENGTH: 27 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear	
	(ii)	MOLECULE TYPE: DNA (genomic)	
	(ix)	FEATURE: (A) NAME/KEY: CDS (B) LOCATION: 127	
	(xi)	SEQUENCE DESCRIPTION: SEQ ID NO:2:	
	AAGGGTGG	TT GTCAGTGGAA TTCGATG	27
	(2) INFO	DRMATION FOR SEQ ID NO:3:	
Salvania 2 2 2 2 3 3 3 3 3 3 3 3 3 3 3 3 3 3 3	(i)	SEQUENCE CHARACTERISTICS: (A) LENGTH: 27 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear	
	(ii)) MOLECULE TYPE: DNA (genomic)	
	(ix)) FEATURE: (A) NAME/KEY: CDS (B) LOCATION: 127	
	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:3:	
	GAGGCCA	GTG CGCCTTGGAA TTCCTAC	27
	(2) INF	ORMATION FOR SEQ ID NO:4:	
	(i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 26 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear	
	(ii) MOLECULE TYPE: DNA (genomic)	
	(ix) FEATURE: (A) NAME/KEY: CDS (B) LOCATION: 126	
	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:4:	
	GCGGTGG	AGG AGACGAAGAA TTCAGT	27
	(2) INF	ORMATION FOR SEQ ID NO:5:	

	(i)	SEQUENCE CHARACTERISTICS: (A) LENGTH: 19 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear	
	(ii)	MOLECULE TYPE: DNA (genomic)	
	(ix)	FEATURE: (A) NAME/KEY: CDS (B) LOCATION: 119	
	(xi)	SEQUENCE DESCRIPTION: SEQ ID NO:5:	
	TTCACTTG	TG CCCTGACTT	19
	(2) INFO	RMATION FOR SEQ ID NO:6:	
	(i)	SEQUENCE CHARACTERISTICS: (A) LENGTH: 20 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear	
	(ii)	MOLECULE TYPE: DNA (genomic)	
The first fi	(ix)	FEATURE: (A) NAME/KEY: CDS (B) LOCATION: 120	
ii Lul	(xi)	SEQUENCE DESCRIPTION: SEQ ID NO:6:	
	CTGGAAAC	TT TCCACTTGAT	20
	(2) INFO	RMATION FOR SEQ ID NO:7:	
Harris Harris	(i)	SEQUENCE CHARACTERISTICS: (A) LENGTH: 18 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear	
	(ii)	MOLECULE TYPE: DNA (genomic)	
	(ix)	FEATURE: (A) NAME/KEY: CDS (B) LOCATION: 118	
	(xi)	SEQUENCE DESCRIPTION: SEQ ID NO:7:	
	CCACTGAC	AA CCACCCTT	18

	(2)	INFO	RMATION FOR SEQ 1D NO:8:		
		(i)	SEQUENCE CHARACTERISTICS: (A) LENGTH: 18 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear		
		(ii)	MOLECULE TYPE: DNA (genomic)		
		(ix)	FEATURE: (A) NAME/KEY: CDS (B) LOCATION: 118		
		(xi)	SEQUENCE DESCRIPTION: SEQ ID NO:8:		
	CCA	AGGCG	CA CTGGCCTC	·•	18
	(2)	INFO	RMATION FOR SEQ ID NO:9:		
		(i)	SEQUENCE CHARACTERISTICS: (A) LENGTH: 18 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear	•	
		(ii)	MOLECULE TYPE: DNA (genomic)	,	
And the first of t		(ix)	FEATURE: (A) NAME/KEY: CDS (B) LOCATION: 118		
osi,		(xi)	SEQUENCE DESCRIPTION: SEQ ID NO:9:		
	CAC	AAACA	ATG CACCTCAA		18
	(2)	INFO	RMATION FOR SEQ ID NO:10:		
		(i)	SEQUENCE CHARACTERISTICS: (A) LENGTH: 18 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear		
		(ii)	MOLECULE TYPE: DNA (genomic)		
		(ix)	FEATURE: (A) NAME/KEY: CDS (B) LOCATION: 118		
		(xi)	SEQUENCE DESCRIPTION: SEQ ID NO:10:		
	TCT	CCCAG	GTA CAGGCACA		18
	(2)	INFO	DRMATION FOR SEQ ID NO:11:		

		(i)	SEQUENCE CHARACTERISTICS: (A) LENGTH: 17 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear				
		(ii)	MOLECULE TYPE: DNA (genomic)				
		(ix)	FEATURE: (A) NAME/KEY: CDS (B) LOCATION: 117				
		(xi)	SEQUENCE DESCRIPTION: SEQ ID NO:11:				
	TGCG	CCGG	CC TCTCCCA	17			
	(2)	INFO	RMATION FOR SEQ ID NO:12:				
Total Barre		(i)	SEQUENCE CHARACTERISTICS: (A) LENGTH: 18 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear				
The state of the s		(ii)	MOLECULE TYPE: DNA (genomic)				
The three states and the states of the state		(ix)	FEATURE: (A) NAME/KEY: CDS (B) LOCATION: 118				
		(xi)	SEQUENCE DESCRIPTION: SEQ ID NO:12:				
	CCGGTTCATG GCGCCCAT						
hei Pli	(2)	INFO	RMATION FOR SEQ ID NO:13:				
The limit of the l		(i)	SEQUENCE CHARACTERISTICS: (A) LENGTH: 27 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear				
		(ii)	MOLECULE TYPE: DNA (genomic)				
		(ix)	FEATURE: (A) NAME/KEY: CDS (B) LOCATION: 127				
		(xi)	SEQUENCE DESCRIPTION: SEQ ID NO:13:				
	GTA	GGAAT	TC ACTTGTGCCC TGACTTG	27			
	(2)	INFO	RMATION FOR SEQ ID NO:14:				

	(L) SEQUENCE CHARACTERISTICS: (A) LENGTH: 29 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear	
	(i	i) MOLECULE TYPE: DNA (genomic)	
	(i:	(A) NAME/KEY: CDS (B) LOCATION: 129	
	(x	i) SEQUENCE DESCRIPTION: SEQ ID NO:14:	
	CATCGA	ATTC TGGAAACTTT CCACTTGAT	29
	(2) IN	FORMATION FOR SEQ ID NO:15:	
	(i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 17 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear	
	(i	i) MOLECULE TYPE: DNA (genomic)	
	(i	x) FEATURE: (A) NAME/KEY: CDS (B) LOCATION: 117	
	(x	i) SEQUENCE DESCRIPTION: SEQ ID NO:15:	
	CCTGCA	GTAG TCCCCTG	17
	(2) IN	FORMATION FOR SEQ ID NO:16:	
The state of the s	(i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 18 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear 	
	(i	i) MOLECULE TYPE: DNA (genomic)	
	(i	x) FEATURE: (A) NAME/KEY: CDS (B) LOCATION: 118	
	(x	i) SEQUENCE DESCRIPTION: SEQ ID NO:16:	
	CCTCAA	CAGG ATGTTTTG	18
	(2) IN	FORMATION FOR SEQ ID NO:17:	

18

(i) SEQUENCE CHARACTERISTICS:

(A) NAME/KEY: CDS (B) LOCATION: 1..18

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:17:

(ix) FEATURE:

(A) LENGTH: 18 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear (ii) MOLECULE TYPE: DNA (genomic)

	(2)	INFOR	MATION FOR SEQ ID NO:20:	
		(i)	SEQUENCE CHARACTERISTICS: (A) LENGTH: 17 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear	
		(ii)	MOLECULE TYPE: DNA (genomic)	
		(ix)	FEATURE: (A) NAME/KEY: CDS (B) LOCATION: 117	
		(xi)	SEQUENCE DESCRIPTION: SEQ ID NO:20:	
	GGCC	ATCC	AC AAGCAGT	17
	(2)	INFO	RMATION FOR SEQ ID NO:21:	
The two that the part of the two that the two the two the two the two the two		(i)	SEQUENCE CHARACTERISTICS: (A) LENGTH: 17 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear	
		(ii)	MOLECULE TYPE: DNA (genomic)	
		(ix)	FEATURE: (A) NAME/KEY: CDS (B) LOCATION: 117	
		(xi)	SEQUENCE DESCRIPTION: SEQ ID NO:21:	
	ACA	GCACA	CG ACGGAGG ·	17
in Fi	(2)	INFO	RMATION FOR SEQ ID NO:22:	
Permi		(i)	SEQUENCE CHARACTERISTICS: (A) LENGTH: 17 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear	
		(ii)	MOLECULE TYPE: DNA (genomic)	
		(ix)	FEATURE: (A) NAME/KEY: CDS (B) LOCATION: 117	
		(xi)	SEQUENCE DESCRIPTION: SEQ ID NO:22:	
	CGG	AGGTT	TCT GAGGCGC	17

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	(2)	INFOR	CHATTON FOR SEQ ID NO:25:	
		(i)	SEQUENCE CHARACTERISTICS: (A) LENGTH: 17 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear	
		(ii)	MOLECULE TYPE: DNA (genomic)	
		(ix)	FEATURE: (A) NAME/KEY: CDS (B) LOCATION: 117	
		(xi)	SEQUENCE DESCRIPTION: SEQ ID NO:23:	
	TTGT	[GAGG	CA CTGCCCC	1.7
	(2)	INFO	RMATION FOR SEQ ID NO:24:	
		(i)	SEQUENCE CHARACTERISTICS: (A) LENGTH: 16 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear	
		(ii)	MOLECULE TYPE: DNA (genomic)	
		(ix)	FEATURE: (A) NAME/KEY: CDS (B) LOCATION: 116	
		(xi)	SEQUENCE DESCRIPTION: SEQ ID NO:24:	
	AGG	CGCTT	CC CCCACC	16
	(2)	INFO	RMATION FOR SEQ ID NO:25:	
122		(i)	SEQUENCE CHARACTERISTICS: (A) LENGTH: 16 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear	
		(ii)	MOLECULE TYPE: DNA (genomic)	
		(ix)	FEATURE: (A) NAME/KEY: CDS (B) LOCATION: 116	
		(xi)	SEQUENCE DESCRIPTION: SEQ ID NO:25:	
	CCC	CACCG	TG AGCGCT	16

	(2)	INFOR	RMATION FOR SEQ ID NO:26:	
		(i)	SEQUENCE CHARACTERISTICS: (A) LENGTH: 19 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear	
		(ii)	MOLECULE TYPE: DNA (genomic)	
		(ix)	FEATURE: (A) NAME/KEY: CDS (B) LOCATION: 119	
		(xi)	SEQUENCE DESCRIPTION: SEQ ID NO:26:	
	ATC	CGAGT	GA AAGGAAATT	19
	(2)	INFO	RMATION FOR SEQ ID NO:27:	
		(i)	SEQUENCE CHARACTERISTICS: (A) LENGTH: 19 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear	
		(ii)	MOLECULE TYPE: DNA (genomic)	
Committee of the control of the cont		(ix)	FEATURE: (A) NAME/KEY: CDS (B) LOCATION: 119	
		(xi)	SEQUENCE DESCRIPTION: SEQ ID NO:27:	
lada Jada	TGT	GGAAG'	TC TTTGGATGA	19
	(2)	INFO	RMATION FOR SEQ ID NO:28:	
THE RESERVE		(i)	SEQUENCE CHARACTERISTICS: (A) LENGTH: 18 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear	
		(ii)	MOLECULE TYPE: DNA (genomic)	
		(ix)	FEATURE: (A) NAME/KEY: CDS (B) LOCATION: 118	
		(xi)	SEQUENCE DESCRIPTION: SEQ ID NO:28:	
	TGT		TAT TTGGATGA	18

(2) INFORMATION FOR SEQ ID NO:29:

CCCCACCTTG AGCGCT

	(2)	INFOR	MATION FOR SEQ ID NO:32:	
		(i)	SEQUENCE CHARACTERISTICS: (A) LENGTH: 18 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear	
		(ii)	MOLECULE TYPE: DNA (genomic)	
		(ix)	FEATURE: (A) NAME/KEY: CDS (B) LOCATION: 118	
		(xi)	SEQUENCE DESCRIPTION: SEQ ID NO:32:	
	ACA:	ragta:	IG GTGGTGCC	18
	(2)	INFO	RMATION FOR SEQ ID NO:33:	
		(i)	SEQUENCE CHARACTERISTICS: (A) LENGTH: 18 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear	
		(ii)	MOLECULE TYPE: DNA (genomic)	
Access the second of the secon		(ix)	FEATURE: (A) NAME/KEY: CDS (B) LOCATION: 118	
		(xi)	SEQUENCE DESCRIPTION: SEQ ID NO:33:	
in and	ACA	TAGTI	TT GTGGTGCC	18
	(2)	INFO	RMATION FOR SEQ ID NO:34:	
		(i)	SEQUENCE CHARACTERISTICS: (A) LENGTH: 16 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear	
		(ii)	MOLECULE TYPE: DNA (genomic)	
		(ix)	FEATURE: (A) NAME/KEY: CDS (B) LOCATION: 116	
		(xi	SEQUENCE DESCRIPTION: SEQ ID NO:34:	
	GG1	rgccc'	TGT GAGCCG	16
	(2)) INF	ORMATION FOR SEQ ID NO:35:	

		(i)	SEQUENCE CHARACTERISTICS: (A) LENGTH: 16 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear	
		(ii)	MOLECULE TYPE: DNA (genomic)	
		(ix)	FEATURE: (A) NAME/KEY: CDS (B) LOCATION: 116	
		(xi)	SEQUENCE DESCRIPTION: SEQ ID NO:35:	
	GGTG	CCCCA	AT GAGCCG	16
	(2)	INFOR	RMATION FOR SEQ ID NO:36:	
77-74 6- 6- 815 S.		(i)	SEQUENCE CHARACTERISTICS: (A) LENGTH: 16 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear	
		(ii)	MOLECULE TYPE: DNA (genomic)	
		(ix)	FEATURE: (A) NAME/KEY: CDS (B) LOCATION: 116	
13 15		(xi)	SEQUENCE DESCRIPTION: SEQ ID NO:36:	
ndh:	GGTC	CCCT	CT GAGCCG	16
	(2)	INFO	RMATION FOR SEQ ID NO:37:	
		(i)	SEQUENCE CHARACTERISTICS: (A) LENGTH: 18 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear	
		(ii)	MOLECULE TYPE: DNA (genomic)	
		(ix)	FEATURE: (A) NAME/KEY: CDS (B) LOCATION: 118	
		(xi)	SEQUENCE DESCRIPTION: SEQ ID NO:37:	
	TGG	CTCTG	GAG TGTACCAC	18
	(2)	INFC	ORMATION FOR SEQ ID NO:38:	
		(i)	SEQUENCE CHARACTERISTICS: (A) LENGTH: 18 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear	

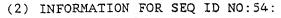
	(ii) MO	OLECULE TYPE: DNA (genomic)	
	` ,	EATURE: (A) NAME/KEY: CDS (B) LOCATION: 118	
	(xi) S	EQUENCE DESCRIPTION: SEQ ID NO:38:	
	CATCCACTGC	AACTACAT	18
	(2) INFORM	ATION FOR SEQ ID NO:39:	
	(i) S	EQUENCE CHARACTERISTICS: (A) LENGTH: 20 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear	
	(ii) M	OLECULE TYPE: DNA (genomic)	
	(ix) F	CEATURE: (A) NAME/KEY: CDS (B) LOCATION: 120	
	(xi) S	EQUENCE DESCRIPTION: SEQ ID NO:39:	0.0
	CTACAACTAA	ATGTGTAACA	20
Milly straig	(2) INFORM	MATION FOR SEQ ID NO:40:	
The state of the s	(i) S	SEQUENCE CHARACTERISTICS: (A) LENGTH: 20 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear	
	(ii) N	MOLECULE TYPE: DNA (genomic)	
Pari.	(ix) 1	FEATURE: (A) NAME/KEY: CDS (B) LOCATION: 120	
	(xi)	SEQUENCE DESCRIPTION: SEQ ID NO:40:	
	CAACTACAT	T TGTAACAGTT	20
	(2) INFOR	MATION FOR SEQ ID NO:41:	
	(i)	SEQUENCE CHARACTERISTICS: (A) LENGTH: 20 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear	
	(ii)	MOLECULE TYPE: DNA (genomic)	

		(1X)	(A) NAME/KEY: CDS (B) LOCATION: 120	
		(xi)	SEQUENCE DESCRIPTION: SEQ ID NO:41:	
	CAAC	TACAT	TA TGTAACAGTT	20
	(2)	INFOR	RMATION FOR SEQ ID NO:42:	
		(i)	SEQUENCE CHARACTERISTICS: (A) LENGTH: 20 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear	
		(ii)	MOLECULE TYPE: DNA (genomic)	
		(ix)	FEATURE: (A) NAME/KEY: CDS (B) LOCATION: 120	
		(xi)	SEQUENCE DESCRIPTION: SEQ ID NO:42:	
	ACTA	ACATGI	TT TAACAGTTCC	20
	(2)	INFO	RMATION FOR SEQ ID NO:43:	
The State		(i)	SEQUENCE CHARACTERISTICS: (A) LENGTH: 20 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear	
		(ii)	MOLECULE TYPE: DNA (genomic)	
		(ix)	FEATURE: (A) NAME/KEY: CDS (B) LOCATION: 120	
		(xi)	SEQUENCE DESCRIPTION: SEQ ID NO:43:	
	ACT	ACATG	TA TAACAGTTCC	20
	(2)	INFO	RMATION FOR SEQ ID NO:44:	
		(i)	SEQUENCE CHARACTERISTICS: (A) LENGTH: 17 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear	
		(ii)	MOLECULE TYPE: DNA (genomic)	
		(ix)	FEATURE: (A) NAME/KEY: CDS (B) LOCATION: 117	

((xi) SEQUENCE DESCRIPTION: SEQ ID NO:44:	
CAGT	ICCTTC ATGGGCG	17
(2)	INFORMATION FOR SEQ ID NO:45:	
	 (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 17 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear 	
1	(ii) MOLECULE TYPE: DNA (genomic)	
	(ix) FEATURE: (A) NAME/KEY: CDS (B) LOCATION: 117	
	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:45:	
GCAT	GGGCGT TATGAAC	17
(2)	INFORMATION FOR SEQ ID NO:46:	
	(i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 17 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear	
	(ii) MOLECULE TYPE: DNA (genomic)	
	(ix) FEATURE: (A) NAME/KEY: CDS (B) LOCATION: 117	
	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:46:	
GCAT	GGGCTG CATGAAC	17
(2)	INFORMATION FOR SEQ ID NO:47:	
	(i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 17 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear	
	(ii) MOLECULE TYPE: DNA (genomic)	
	(ix) FEATURE: (A) NAME/KEY: CDS (B) LOCATION: 117	
	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:47:	
CCAT	COCCCCA CATCAAC	17

	(2)	INFOR	MATION FOR SEQ ID NO:48:		
		(i)	SEQUENCE CHARACTERISTICS: (A) LENGTH: 17 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear		
		(ii)	MOLECULE TYPE: DNA (genomic)		
		(ix)	FEATURE: (A) NAME/KEY: CDS (B) LOCATION: 117		
		(xi)	SEQUENCE DESCRIPTION: SEQ ID NO:48:		
	GGC	GGCTT	GA ACCGGAG		17
	(2)	INFO	RMATION FOR SEQ ID NO:49:		
		(i)	SEQUENCE CHARACTERISTICS: (A) LENGTH: 18 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear	• • • • • • • • • • • • • • • • • • • •	
		(ii)	MOLECULE TYPE: DNA (genomic)		
		(ix)	FEATURE: (A) NAME/KEY: CDS (B) LOCATION: 118		
		(xi)	SEQUENCE DESCRIPTION: SEQ ID NO:49:		
rj Hj	CAT	GAACC	TG AGGCCCAT		18
ASTA	(2)	INFO	RMATION FOR SEQ ID NO:50:		
		(i)	SEQUENCE CHARACTERISTICS: (A) LENGTH: 18 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear		
		(ii)	MOLECULE TYPE: DNA (genomic)		
		(ix)	FEATURE: (A) NAME/KEY: CDS (B) LOCATION: 118		
		(xi)	SEQUENCE DESCRIPTION: SEQ ID NO:50:		
	GCA	ATGAAC	CTG GAGGCCCA	-	18

	(2)	INFOR	MATION FOR SEQ ID NO:51:	
		(i)	SEQUENCE CHARACTERISTICS: (A) LENGTH: 18 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear	
		(ii)	MOLECULE TYPE: DNA (genomic)	
		(ix)	FEATURE: (A) NAME/KEY: CDS (B) LOCATION: 118	
		(xi)	SEQUENCE DESCRIPTION: SEQ ID NO:51:	
	GCAT	CGAACC	CA GAGGCCCA	18
	(2)	INFO	RMATION FOR SEQ ID NO:52:	
		(i)	SEQUENCE CHARACTERISTICS: (A) LENGTH: 18 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear	
gon gon gon gon gon		(ii)	MOLECULE TYPE: DNA (genomic)	
72		(ix)	FEATURE: (A) NAME/KEY: CDS (B) LOCATION: 118	
		(xi)	SEQUENCE DESCRIPTION: SEQ ID NO:52:	
	AAC	CGGAG'	IC CCATCCTC	18
	(2)	INFO	RMATION FOR SEQ ID NO:53:	
The state of the s		(i)	SEQUENCE CHARACTERISTICS: (A) LENGTH: 16 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear	
		(ii)	MOLECULE TYPE: DNA (genomic)	
		(ix)	FEATURE: (A) NAME/KEY: CDS (B) LOCATION: 116	
		(xi)	SEQUENCE DESCRIPTION: SEQ ID NO:53:	
	GAA	.ccggg	GG CCCATC	16



- (i) SEQUENCE CHARACTERISTICS:

 (A) LENGTH: 17 base pairs

 (B) TYPE: nucleic acid

 (C) STRANDEDNESS: single

 (D) TOPOLOGY: linear
- (ii) MOLECULE TYPE: DNA (genomic)
- (ix) FEATURE:
 (A) NAME/KEY: CDS
 (B) LOCATION: 1..17

		(xi)	SEQUENCE DESCRIPTION: SEQ ID NO:54:	
	GAGG	CCCAA	AC CTCACCA	17
	(2)	INFOR	RMATION FOR SEQ ID NO:55:	
		(i)	SEQUENCE CHARACTERISTICS: (A) LENGTH: 17 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear	
		(ii)	MOLECULE TYPE: DNA (genomic)	
		(ix)	FEATURE: (A) NAME/KEY: CDS (B) LOCATION: 117	
		(xi)	SEQUENCE DESCRIPTION: SEQ ID NO:55:	
	CATO	CACAC	CG GAAGACT	17
	(2)	INFO	RMATION FOR SEQ ID NO:56:	
		(i)	SEQUENCE CHARACTERISTICS: (A) LENGTH: 18 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear	
2 - 2 - 2 - 2 - 2 - 2 - 2 - 2 - 2 - 2 -		(ii)	MOLECULE TYPE: DNA (genomic)	
Sant Care Hr. Tark Hr.		(ix)	FEATURE: (A) NAME/KEY: CDS (B) LOCATION: 118	
		(xi)	SEQUENCE DESCRIPTION: SEQ ID NO:56:	
THE PERSON NAMED IN	TCTA	ACTGG	AA CGGAACAG	18
	(2)	INFO	RMATION FOR SEQ ID NO:57:	
		(i)	SEQUENCE CHARACTERISTICS: (A) LENGTH: 18 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear	
		(ii)	MOLECULE TYPE: DNA (genomic)	
		(ix)	FEATURE: (A) NAME/KEY: CDS (B) LOCATION: 118	
		(xi)	SEQUENCE DESCRIPTION: SEQ ID NO:57:	
	TTG	AGGTG	CA TGTTTGTG	18

(2)	INFOR	RMATION FOR SEQ ID NO:58:	
	(i)	SEQUENCE CHARACTERISTICS: (A) LENGTH: 18 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear	
	(ii)	MOLECULE TYPE: DNA (genomic)	
	(ix)	FEATURE: (A) NAME/KEY: CDS (B) LOCATION: 118	
	(xi)	SEQUENCE DESCRIPTION: SEQ ID NO:58:	
TTGA	AGGTG	GG TGTTTGTG	18
(2)	INFO	RMATION FOR SEQ ID NO:59:	
	(i)	SEQUENCE CHARACTERISTICS: (A) LENGTH: 17 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear	
	(ii)	MOLECULE TYPE: DNA (genomic)	
	(ix)	FEATURE: (A) NAME/KEY: CDS (B) LOCATION: 117	
	(xi)	SEQUENCE DESCRIPTION: SEQ ID NO:59:	
GCT	GTTTA'	IG CCTGCCT	17
(2)	INFO	RMATION FOR SEQ ID NO:60:	
	(i)	SEQUENCE CHARACTERISTICS: (A) LENGTH: 18 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear	
	(ii)	MOLECULE TYPE: DNA (genomic)	
	(ix)	FEATURE: (A) NAME/KEY: CDS (B) LOCATION: 118	
	(xi)	SEQUENCE DESCRIPTION: SEQ ID NO:60:	
TGC	CTGTA	CT GGGAGAGA	18

	(2)	INFOR	MATION FOR SEQ ID NO:61:	
		(i)	SEQUENCE CHARACTERISTICS: (A) LENGTH: 18 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear	
		(ii)	MOLECULE TYPE: DNA (genomic)	
A the second sec		(ix)	FEATURE: (A) NAME/KEY: CDS (B) LOCATION: 118	
		(xi)	SEQUENCE DESCRIPTION: SEQ ID NO:61:	
	TGC	CTGTT	CT GGGAGAGA	18
	(2)	INFO	RMATION FOR SEQ ID NO:62:	
		(i)	SEQUENCE CHARACTERISTICS: (A) LENGTH: 18 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear	
		(ii)	MOLECULE TYPE: DNA (genomic)	
		(ix)	FEATURE: (A) NAME/KEY: CDS (B) LOCATION: 118	
		(xi)	SEQUENCE DESCRIPTION: SEQ ID NO:62:	
pa Fil	TGC	CTGTC	GT GGGAGAGA	18
	(2)	INFO	RMATION FOR SEQ ID NO:63:	
		(i)	SEQUENCE CHARACTERISTICS: (A) LENGTH: 17 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear	
		(ii)	MOLECULE TYPE: DNA (genomic)	
		(ix)	FEATURE: (A) NAME/KEY: CDS (B) LOCATION: 117	
		(xi)	SEQUENCE DESCRIPTION: SEQ ID NO:63:	
	CTG	TCCTG	AG AGAGACC	17

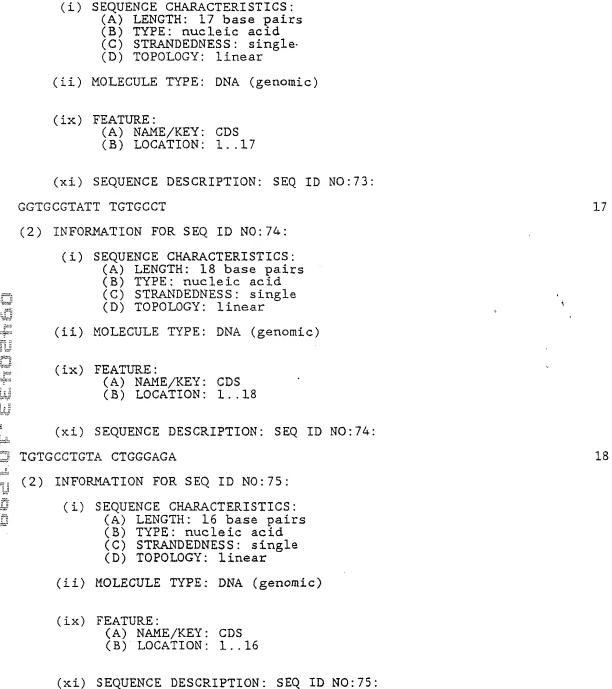
	(2)	INFOR	MATION FOR SEQ ID NO:64:	
		(i)	SEQUENCE CHARACTERISTICS: (A) LENGTH: 17 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear	
		(ii)	MOLECULE TYPE: DNA (genomic)	
		(ix)	FEATURE: (A) NAME/KEY: CDS (B) LOCATION: 117	
		(xi)	SEQUENCE DESCRIPTION: SEQ ID NO:64:	
	CTGT	CCTGC	GG GAGAGAC	17
	(2)	INFOR	RMATION FOR SEQ ID NO:65:	
		(i)	SEQUENCE CHARACTERISTICS: (A) LENGTH: 17 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear	
		(ii)	MOLECULE TYPE: DNA (genomic)	
i#		(ix)	FEATURE: (A) NAME/KEY: CDS (B) LOCATION: 117	
		(xi)	SEQUENCE DESCRIPTION: SEQ ID NO:65:	
TJ.	TGG	GAGAC	AC CGGCGCA	17
	(2)	INFO	RMATION FOR SEQ ID NO:66:	
		(i)	SEQUENCE CHARACTERISTICS: (A) LENGTH: 17 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear	
		(ii)	MOLECULE TYPE: DNA (genomic)	
		(ix)	FEATURE: (A) NAME/KEY: CDS (B) LOCATION: 117	
		(xi)	SEQUENCE DESCRIPTION: SEQ ID NO:66:	4 =
	TG	GGAGA	GAG CGGCGCA .	17
	(2) INFO	DRMATION FOR SEQ ID NO:67:	

		(i)	SEQUENCE CHARACTERISTICS: (A) LENGTH: 18 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear				
		(ii)	MOLECULE TYPE: DNA (genomic)				
		(ix)	FEATURE: (A) NAME/KEY: CDS (B) LOCATION: 118				
		(xi)	SEQUENCE DESCRIPTION: SEQ ID NO:67:				
	GCGC	ACAAA	AG GAAGAGAA	18			
	(2)	INFO	RMATION FOR SEQ ID NO:68:				
		(i)	SEQUENCE CHARACTERISTICS: (A) LENGTH: 19 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear				
		(ii)	MOLECULE TYPE: DNA (genomic)				
		(ix)	FEATURE: (A) NAME/KEY: CDS (B) LOCATION: 119				
		(xi)	SEQUENCE DESCRIPTION: SEQ ID NO:68:				
	GCAG	GCACAGAGAA AGAGAATCT					
	(2)	INFO	RMATION FOR SEQ ID NO:69:				
		(i)	SEQUENCE CHARACTERISTICS: (A) LENGTH: 17 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear				
		(ii)	MOLECULE TYPE: DNA (genomic)				
		(ix)	FEATURE: (A) NAME/KEY: CDS (B) LOCATION: 117				
		(xi)	SEQUENCE DESCRIPTION: SEQ ID NO:69:				
	GAA.	AGGGT	AG CCTCACC	17			
	(2)	INFO	RMATION FOR SEQ ID NO:70:				



	(i)	SEQUENCE CHARACTERISTICS: (A) LENGTH: 17 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear	
	(ii)	MOLECULE TYPE: DNA (genomic)	
	(ix)	FEATURE: (A) NAME/KEY: CDS (B) LOCATION: 117	
	(xi)	SEQUENCE DESCRIPTION: SEQ ID NO:70:	
GA	GCCTCC	CC ACGAGCT	17
(2) INFO	RMATION FOR SEQ ID NO:71:	
Market Ma Market Market Market Ma Ma Ma Ma Ma Ma Ma Ma Ma Ma Ma Ma Ma	(i)	SEQUENCE CHARACTERISTICS: (A) LENGTH: 17 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear	
	(ii)	MOLECULE TYPE: DNA (genomic)	
	(ix)	FEATURE: (A) NAME/KEY: CDS (B) LOCATION: 117	
	(xi)	SEQUENCE DESCRIPTION: SEQ ID NO:71:	
i ci	CACCAC	CTA GCTGCCC	17
	2) INFO	RMATION FOR SEQ ID NO:72:	
	(i)	SEQUENCE CHARACTERISTICS: (A) LENGTH: 17 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear	
	(ii)) MOLECULE TYPE: DNA (genomic)	
	(ix)	FEATURE: (A) NAME/KEY: CDS (B) LOCATION: 117	
	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:72:	
С	TCACCA	CGC ACTGCCC	17
(2) INF	ORMATION FOR SEQ ID NO:73:	

TGGGAGACAC CGGCGC





(2) INFORMATION FOR SEQ ID NO:79:

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1			
	-	3	
4		₹	

(2)	INFOR	RMATION FOR SEQ ID NO:76:	
	(i)	SEQUENCE CHARACTERISTICS: (A) LENGTH: 31 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear	
	(ii)	MOLECULE TYPE: DNA (genomic)	
	(ix)	FEATURE: (A) NAME/KEY: CDS (B) LOCATION: 131	
	(xi)	SEQUENCE DESCRIPTION: SEQ ID NO:76:	
CAUC	CAUCAU	UC AUTTCACTTG TGCCCTGACT T	31
(2)	INFO	RMATION FOR SEQ ID NO:77:	
s in hink their	(i)	SEQUENCE CHARACTERISTICS: (A) LENGTH: 32 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear	
	(ii)	MOLECULE TYPE: DNA (genomic)	
	(ix)	FEATURE: (A) NAME/KEY: CDS (B) LOCATION: 132	
	(xi)	SEQUENCE DESCRIPTION: SEQ ID NO:77:	
CAU	CAUCU	AC UACTGGAAAC TTTCCACTTG AT	32
(2)	INFO	RMATION FOR SEQ ID NO:78:	
	(i)	SEQUENCE CHARACTERISTICS: (A) LENGTH: 30 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear	
	(ii)	MOLECULE TYPE: DNA (genomic)	
	(ix)	FEATURE: (A) NAME/KEY: CDS (B) LOCATION: 130	
	(xi)	SEQUENCE DESCRIPTION: SEQ ID NO:78:	•
CUA	CUACU	AC UACCACTGAC AACCACCCTT	30





	(i)	SEQUENCE CHARACTERISTICS: (A) LENGTH: 30 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear	
	(ii)	MOLECULE TYPE: DNA (genomic)	
	(ix)	FEATURE: (A) NAME/KEY: CDS (B) LOCATION: 130	
	(xi)	SEQUENCE DESCRIPTION: SEQ ID NO:79:	
	CAUCAUCA	UC AUCCAAGGCG CACTGGCCTC	30
	(2) INFO	RMATION FOR SEQ ID NO:80:	
	(i)	SEQUENCE CHARACTERISTICS: (A) LENGTH: 17 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear	
	(ii)	MOLECULE TYPE: DNA (genomic)	
	(ix)	FEATURE: (A) NAME/KEY: CDS (B) LOCATION: 117	
is Jack Jack	(xi)	SEQUENCE DESCRIPTION: SEQ ID NO:80:	
L.	TGGGAGAG	GC CGGCGCA	17
	(2) INFO	RMATION FOR SEQ ID NO:81:	
	(i)	SEQUENCE CHARACTERISTICS: (A) LENGTH: 18 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear	
	(ii)	MOLECULE TYPE: DNA (genomic)	
	(ix)	FEATURE: (A) NAME/KEY: CDS (B) LOCATION: 118	
	(xí)	SEQUENCE DESCRIPTION: SEQ ID NO:81:	
	ATGGGCGC	CA TGAACCGG	18



- (i) SEQUENCE CHARACTERISTICS:

 (A) LENGTH: 18 base pairs

 (B) TYPE: nucleic acid

 (C) STRANDEDNESS: single

 (D) TOPOLOGY: linear
- (ii) MOLECULE TYPE: DNA (genomic)
- (ix) FEATURE:
 - (A) NAME/KEY: CDS (B) LOCATION: 1..18
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO:82:

TGTGCCTGTA CTGGGAGA